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CORRIGENDUM

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The authors wish to correct the reported genome sequence of the bovine astrovirus CH15 (BoAstV-CH15) strain (GenBank accession number KT956903.1). Resequencing of the entire viral genome by two additional runs of Sanger sequencing of RT-PCR amplicons in addition to 5' and 3' RACE showed previously unrecognized additional 171 nt at the 5' end of the viral genome and a 9 nt insertion in open reading frame 1a. The GenBank database entry has been updated accordingly (GenBank accession number KT956903.2).

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